

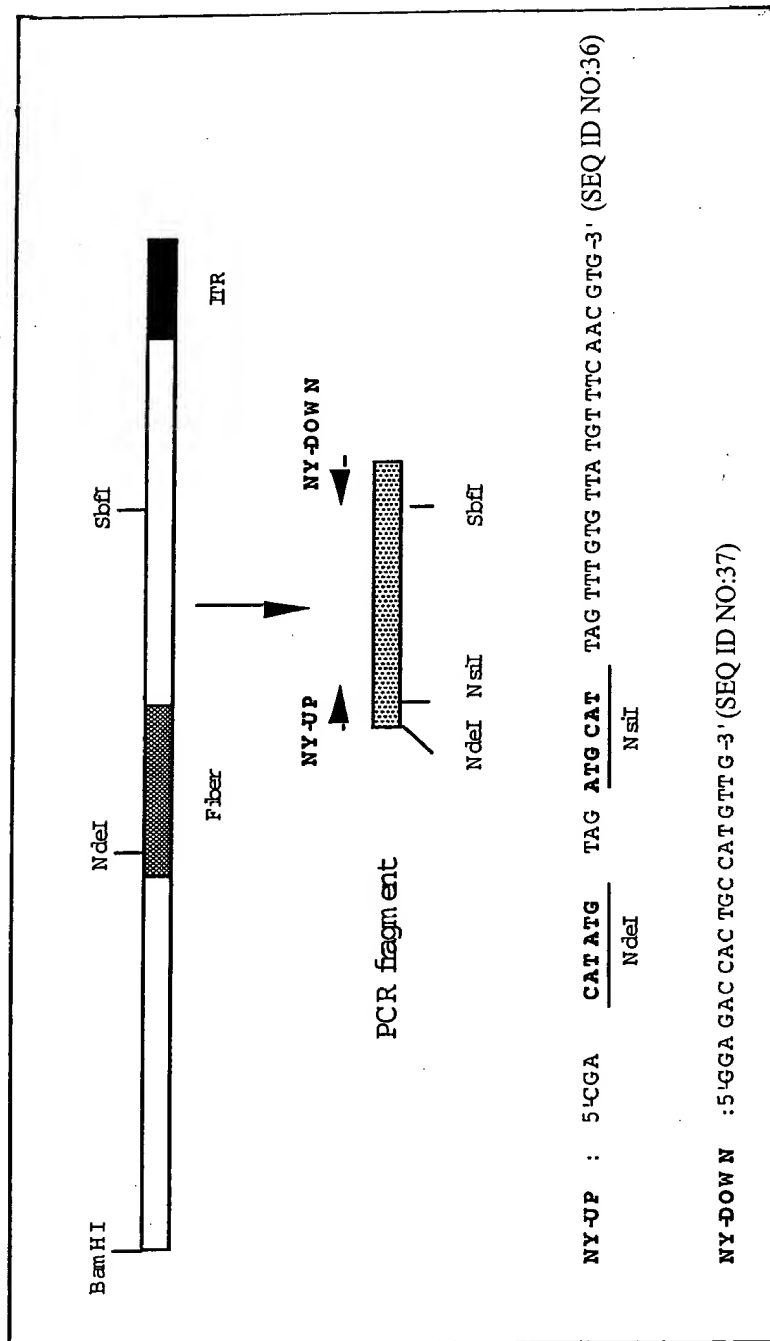


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Figure 2



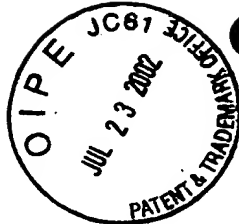


Figure 4: Sequence of Ad5/fib16 chimeric fiber

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TCTAACTCTTAAATGTGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTTAAAGTTGGAAGC
AGTCTTACAGTAGATACTATCGATGGGTCTTTGGAGGAAAATATAACTGCCGAAGCGCCACTCACTA
AAACTAACCCTCCATAGGTTTATTAATAGGATCTGGCTTGCAAACAAAGGATGATAAACTTTGTTT
ATCGCTGGGAGATGGGTTGGTAACAAAGGATGATAAACTATGTTTATCGCTGGGAGATGGGTTAATA
ACAAAAAATGATGTACTATGTGCCAACTAGGACATGGCCTTGTGTTTGACTCTTCCAATGCTATCA
CCATAGAAAACAACACCTTGTGGACAGGCGCAAAACCAAGCGCCAACCTGTGTAATTAAAGAGGGAGA
AGATTCCCCAGACTGTAAGCTCACTTTAGTTCTAGTGAAGAATGGAGGACTGATAAATGGATACATA
ACATTAATGGGAGCCTCAGAATATACTAACACCTTGTTTAAAAACAATCAAGTTACAATCGATGTAA
ACCTCGCATTTGATAATACTGGCCAAATTATTACTTACCTATCATCCCTTAAAAGTAACCTGAACCTT
TAAAGACAACCAAAACATGGCTACTGGAACCATAACCAGTGCCAAAGGCTTCATGCCCAGCACCACC
GCCTATCCATTTATAACATACGCCACTGAGACCCTAAATGAAGATTACATTTATGGAGAGTGTTACT
ACAAATCTACCAATGGAACCTCTTTTCCACTAAAAGTTACTGTCACACTAAACAGACGTATGTTAGC
TTCTGGAATGGCCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGGAACTACC
GAAGTCACTCTCATTACCTCCCCCTTCTTTTTTCTTATATCAGAGAAGATGACTGAATGCATTAG
(SEQ ID NO:32)

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FIGURE 7A

Alignment Report of alignment.DNA, using Clustal method with Weighted residue weight table.
Wednesday, March 31, 1999 16:13

1 ATGGC - CAAACGAGCTCGGCTAAGCAGCTCCTTCAATCCGGTCTACCCCT Fiber 16 Genbank (SEQ ID NO:33)
1 ATG[AAG]C[GCG]C[A]AG[A]C[G]T[G]A[A-GAT]A CCTTCAA[C]C[C]G T[G]TA[T]C C[A]T Ad5/16 Fiber DNA (SEQ ID NO:38)

50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTTATAAAACCCCTGGTTTTCATT Fiber 16 Genbank
50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTTATAAAACCCCTGGTTTTCATT Ad5/16 Fiber DNA

100 TCCTCAAATGGTTTTTGCACAAAGCCAGATGGAGTTCTAACTCTTAAATG Fiber 16 Genbank
100 TCCTCAAATGGTTTTTGCACAAAGCCAGATGGAGTTCTAACTCTTAAATG Ad5/16 Fiber DNA

150 TGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTTAAAGTTGGAA Fiber 16 Genbank
150 TGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTTAAAGTTGGAA Ad5/16 Fiber DNA

200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTTGGAGGAAAAATATAACT Fiber 16 Genbank
200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTTGGAGGAAAAATATAACT Ad5/16 Fiber DNA

250 GCCGCAGCGCCACTCACTAAAACCTAACCACCTCCATAGGTTTATTAATAGG Fiber 16 Genbank
250 GCCG[A]AGCGCCACTCACTAAAACCTAACCACCTCCATAGGTTTATTAATAGG Ad5/16 Fiber DNA

300 ATCTGGCTTGCACAAACAAAGGATGATAAACTTTGTTTATCGCTGGGAGATG Fiber 16 Genbank
300 ATCTGGCTTGCACAAACAAAGGATGATAAACTTTGTTTATCGCTGGGAGATG Ad5/16 Fiber DNA

350 GGTTGGTAACAAAGGATGATAAACTATGTTTATCGCTGGGAGATGGGTTA Fiber 16 Genbank
350 GGTTGGTAACAAAGGATGATAAACTATGTTTATCGCTGGGAGATGGGTTA Ad5/16 Fiber DNA

400 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTGTT Fiber 16 Genbank
406 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTGTT Ad5/16 Fiber DNA

450 TGACTCTTCCAATGCTATCACCATAGAAAAACAACACCTTGTGGACAGGCG Fiber 16 Genbank
450 TGACTCTTCCAATGCTATCACCATAGAAAAACAACACCTTGTGGACAGGCG Ad5/16 Fiber DNA

500 CAAAACCAAGCGCCAACTGTGTAATTAAAGAGGGAGAAAGATTCCCCAGAC Fiber 16 Genbank
500 CAAAACCAAGCGCCAACTGTGTAATTAAAGAGGGAGAAAGATTCCCCAGAC Ad5/16 Fiber DNA

550 TGTAAGCTCACTTTAGTTCTAGTGAAGAATGGAGGACTGATAAATGGATA Fiber 16 Genbank
550 TGTAAGCTCACTTTAGTTCTAGTGAAGAATGGAGGACTGATAAATGGATA Ad5/16 Fiber DNA

600 CATAACATTAATGGGAGCCTCAGAATATACTAACACCTTGTTTAAAAACA Fiber 16 Genbank
600 CATAACATTAATGGGAGCCTCAGAATATACTAACACCTTGTTTAAAAACA Ad5/16 Fiber DNA

650 ATCAAGTTACAATCGATGTAAACCTCGCATTTTGATAAATACTGGCCAAATT Fiber 16 Genbank
650 ATCAAGTTACAATCGATGTAAACCTCGCATTTTGATAAATACTGGCCAAATT Ad5/16 Fiber DNA

700 ATTACTTACCTATCATCCCTTAAAAGTAACCTGAACCTTAAAGACAACCA Fiber 16 Genbank
700 ATTACTTACCTATCATCCCTTAAAAGTAACCTGAACCTTAAAGACAACCA Ad5/16 Fiber DNA

750 AAACATGGCTACTGGAACCATAAACAGTGCCAAAGGCTTCATGCCCAGCA Fiber 16 Genbank
750 AAACATGGCTACTGGAACCATAAACAGTGCCAAAGGCTTCATGCCCAGCA Ad5/16 Fiber DNA

800 CCACCGCCTATCCATTTATAACATACGCCACTGAGACCCTAAATGAAGAT Fiber 16 Genbank
800 CCACCGCCTATCCATTTATAACATACGCCACTGAGACCCTAAATGAAGAT Ad5/16 Fiber DNA

850 TACATTTATGGAGAGTGTACTACAAATCTACCAATGGAACCTCTCTTTCC Fiber 16 Genbank
850 TACATTTATGGAGAGTGTACTACAAATCTACCAATGGAACCTCTCTTTCC Ad5/16 Fiber DNA

900 ACTAAAAGTTACTGTCACTAAACAGACGTATGTTAGCTTCTTGGAATGG Fiber 16 Genbank
900 ACTAAAAGTTACTGTCACTAAACAGACGTATGTTAGCTTCTTGGAATGG Ad5/16 Fiber DNA

950 CCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGGAA Fiber 16 Genbank
950 CCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGGAA Ad5/16 Fiber DNA

1000 ACTACCGAAGTCACTCTCATTAACCTCCCCCTTCTTTTTTTCTTATATCAG Fiber 16 Genbank
1000 ACTACCGAAGTCACTCTCATTAACCTCCCCCTTCTTTTTTTCTTATATCAG Ad5/16 Fiber DNA

1050 AGAAGATGACTGA Fiber 16 Genbank
1050 AGAAGATGACTGAATGCATTAGTTTGTGTTATGTTTCAACGTGTTTATTT Ad5/16 Fiber DNA

1062 Fiber 16 Genbank
1100 TCAATTG Ad5/16 Fiber DNA

Decoration 'Decoration #1': Box residues that differ from Fiber 16 Genbank.

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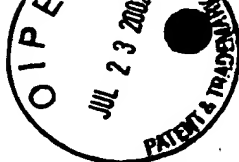


FIGURE 7B

Alignment Report of alignment.prot, using Clustal method with PAM250 residue weight table.
Wednesday, March 31, 1999 16:19

Page 1

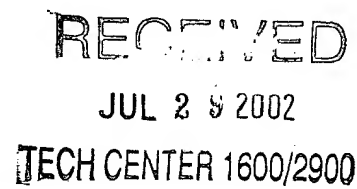
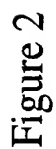
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1	M - K R A R P S E D T F N P V Y P Y E D E S S S Q H P F I N	chimeric Ad5/Fib16	(SEQ ID NO:34)
30	P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G	genbank Fiber 16	
30	P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G	chimeric Ad5/Fib16	
60	P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T	genbank Fiber 16	
60	P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T	chimeric Ad5/Fib16	
90	K T N H S I G L L I G S G L Q T K D D K L C L S L G D G L V	genbank Fiber 16	
90	K T N H S I G L L I G S G L Q T K D D K L C L S L E D G L V	chimeric Ad5/Fib16	
120	T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V	genbank Fiber 16	
120	T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V	chimeric Ad5/Fib16	
150	F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E	genbank Fiber 16	
150	F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E	chimeric Ad5/Fib16	
180	D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y	genbank Fiber 16	
180	D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y	chimeric Ad5/Fib16	
210	T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S	genbank Fiber 16	
210	T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S	chimeric Ad5/Fib16	
240	L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A	genbank Fiber 16	
240	L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A	chimeric Ad5/Fib16	
270	Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F	genbank Fiber 16	
270	Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F	chimeric Ad5/Fib16	
300	P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E	genbank Fiber 16	
300	P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E	chimeric Ad5/Fib16	
330	E A P E T T E V T L I T S P F F F S Y I R E D D	genbank Fiber 16	
330	E A P E T T E V T L I T S P F F F S Y I R E D D	chimeric Ad5/Fib16	

Decoration 'Decoration #1': Box residues that differ from genbank Fiber 16 .

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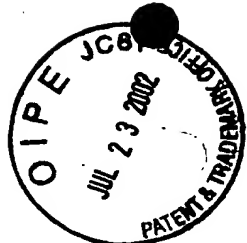


Figure 4: Sequence of Ad5/fib16 chimeric fiber

ATGAAGCGCGCAAGACCGTCTGAAGATACCTTCAACCCCGTGTATCCATATGAAGATGAAAGCAGCT
CACAACACCCCTTTATAAACCCCTGGTTTCATTTCCCTCAAATGGTTTTGCACAAAGCCCAGATGGAGT
TCTAACTCTTAAATGTGTAAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTAAAGTTGGAAGC
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AAACTAACCCTCCATAGGTTTATTAATAGGATCTGGCTTGCAAACAAAGGATGATAAACTTTGTTT
ATCGCTGGGAGATGGGTGGTAACAAAGGATGATAAACTATGTTTATCGCTGGGAGATGGGTAAATA
ACAAAAAATGATGTACTATGTGCCAACTAGGACATGGCCTTGTGTTTGACTCTTCCAATGCTATCA
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TAAAGACAACCAAAACATGGCTACTGGAACCATAACCAGTGCCAAAGGCTTCATGCCCAGCACCACC
GCCTATCCATTTATAACATACGCCACTGAGACCCATAATGAAGATTACATTTATGGAGAGTGTTACT
ACAAATCTACCAATGGAACCTCTTTTCCACTAAAAGTTACTGTCACTAAACAGACGTATGTTAGC
TTCTGGAATGGCCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGGAACTACC
GAAGTCACTCTCATTACCTCCCCCTTCTTTTTTTCTTATATCAGAGAAGATGACTGAATGCATTAG

(SEQ ID NO: 32)

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FIGURE 7A

Alignment Report of alignment.DNA, using Clustal method with Weighted residue weight table.
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Page 1

1 ATGGC - CAAACGAGCTCGGCTAAGCAGCTCCTTCAATCCGGTCTACCCCT Fiber 16 Genbank (Seq ID NO: 38)
1 ATG(AAG)C(GCG)C(AAG)A(C)G(T)C(T)G(A)A(GATA)CCTTCAA(C)C(C)G(T)G(T)A(T)C(C)A(T) Ad5/16 Fiber DNA
50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTTATAAACCCTGGTTTTCATT Fiber 16 Genbank
50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTTATAAACCCTGGTTTTCATT Ad5/16 Fiber DNA
100 TCCTCAAATGGTTTTTGCACAAAGCCCAGATGGAGTTCTAACTCTTAAATG Fiber 16 Genbank
100 TCCTCAAATGGTTTTTGCACAAAGCCCAGATGGAGTTCTAACTCTTAAATG Ad5/16 Fiber DNA
150 TGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTTAAAGTTGGAA Fiber 16 Genbank
150 TGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTTAAAGTTGGAA Ad5/16 Fiber DNA
200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTTGGAGGAAAATATAACT Fiber 16 Genbank
200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTTGGAGGAAAATATAACT Ad5/16 Fiber DNA
250 GCCGCAGCGCCACTCACTAAAACCTAACCACTCCATAGGTTTATTAAATAGG Fiber 16 Genbank
250 GCCG(A)AGCGCCACTCACTAAAACCTAACCACTCCATAGGTTTATTAAATAGG Ad5/16 Fiber DNA
300 ATCTGGCTTGCAAACAAAGGATGATAAACTTTGTTTTATCGCTGGGAGATG Fiber 16 Genbank
300 ATCTGGCTTGCAAACAAAGGATGATAAACTTTGTTTTATCGCTGGGAGATG Ad5/16 Fiber DNA
350 GGTGGTTAAACAAAGGATGATAAACTATGTTTTATCGCTGGGAGATGGGTTA Fiber 16 Genbank
350 GGTGGTTAAACAAAGGATGATAAACTATGTTTTATCGCTGGGAGATGGGTTA Ad5/16 Fiber DNA
400 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTTGTT Fiber 16 Genbank
400 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTTGTT Ad5/16 Fiber DNA
450 TGACTCTTCCAATGCTATCACCATAGAAAACAACACCTTGTTGGACAGGCG Fiber 16 Genbank
450 TGACTCTTCCAATGCTATCACCATAGAAAACAACACCTTGTTGGACAGGCG Ad5/16 Fiber DNA
500 CAAAACCAAGCGCCAACTGTGTAAATTAAAGAGGGAGAAAGATTCCCCAGAC Fiber 16 Genbank
500 CAAAACCAAGCGCCAACTGTGTAAATTAAAGAGGGAGAAAGATTCCCCAGAC Ad5/16 Fiber DNA
550 TGTAAAGCTCACTTTTAGTTCTAGTGAAAGAAATGGAGGACTGATAAATGGATA Fiber 16 Genbank
550 TGTAAAGCTCACTTTTAGTTCTAGTGAAAGAAATGGAGGACTGATAAATGGATA Ad5/16 Fiber DNA
600 CATAACATTAATGGGAGCCTCAGAAATATACTAACACCTTGTTTAAAAACA Fiber 16 Genbank
600 CATAACATTAATGGGAGCCTCAGAAATATACTAACACCTTGTTTAAAAACA Ad5/16 Fiber DNA
650 ATCAAGTTACAATCGATGTAAACCTCGCATTTTGATAAATACTGGCCAAATT Fiber 16 Genbank
650 ATCAAGTTACAATCGATGTAAACCTCGCATTTTGATAAATACTGGCCAAATT Ad5/16 Fiber DNA
700 ATTACTTACCTATCATCCCTTAAAAGTAACCTGAACTTTTAAAGACAACCA Fiber 16 Genbank
700 ATTACTTACCTATCATCCCTTAAAAGTAACCTGAACTTTTAAAGACAACCA Ad5/16 Fiber DNA
750 AAACATGGCTACTGGAACCATAAACCAAGTGCCAAAGGCTTCATGCCCAGCA Fiber 16 Genbank
750 AAACATGGCTACTGGAACCATAAACCAAGTGCCAAAGGCTTCATGCCCAGCA Ad5/16 Fiber DNA
800 CCACCGCCTATCCATTTTATAACATACGCCACTGAGACCCTAAATGAAGAT Fiber 16 Genbank
800 CCACCGCCTATCCATTTTATAACATACGCCACTGAGACCCTAAATGAAGAT Ad5/16 Fiber DNA
850 TACATTTTATGGAGAGTGTTACTACAAATCTACCAATGGAACCTCTCTTTCC Fiber 16 Genbank
850 TACATTTTATGGAGAGTGTTACTACAAATCTACCAATGGAACCTCTCTTTCC Ad5/16 Fiber DNA
900 ACTAAAAGTTACTGTCACACTAAACAGACGTAATGTTAGCTTCTGGAATGG Fiber 16 Genbank
900 ACTAAAAGTTACTGTCACACTAAACAGACGTAATGTTAGCTTCTGGAATGG Ad5/16 Fiber DNA
950 CCTATGCTATGAATTTTTCATGGTCTCTAAATGTCAGAGGAAGCCCCGGAA Fiber 16 Genbank
950 CCTATGCTATGAATTTTTCATGGTCTCTAAATGTCAGAGGAAGCCCCGGAA Ad5/16 Fiber DNA
1000 ACTACCGAAGTCACTCTCATTAACCTCCCCCTTCTTTTCTTTTCTTATATCAG Fiber 16 Genbank
1000 ACTACCGAAGTCACTCTCATTAACCTCCCCCTTCTTTTCTTTTCTTATATCAG Ad5/16 Fiber DNA
1050 AGAAGATGACTGA Fiber 16 Genbank
1050 AGAAGATGACTGAATGCATTAGTTTGTGTTATGTTTCAACGTGTTTATTT Ad5/16 Fiber DNA
1062 Fiber 16 Genbank
1100 TCAATTG Ad5/16 Fiber DNA

Decoration 'Decoration #1': Box residues that differ from Fiber 16 Genbank.

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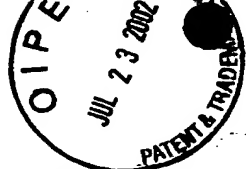


FIGURE 7B

Alignment Report of alignment.prot, using Clustal method with PAM250 residue weight table.
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Page 1

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1  M A K R A R L S S - S F N P V Y P Y E D E S S S Q H P F I N  genbank Fiber 16 (SEQ ID NO: 35)
1  M - K R A R P S E D T F N P V Y P Y E D E S S S Q H P F I N  chimeric Ad5/Fib16 (SEQ ID NO: 34)

30  P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G  genbank Fiber 16
30  P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G  chimeric Ad5/Fib16

60  P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T  genbank Fiber 16
60  P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T  chimeric Ad5/Fib16

90  K T N H S I G L L I G S G L Q T K D D K L C L S L G D G L V  genbank Fiber 16
90  K T N H S I G L L I G S G L Q T K D D K L C L S L E D G L V  chimeric Ad5/Fib16

120 T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V  genbank Fiber 16
120 T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V  chimeric Ad5/Fib16

150 F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E  genbank Fiber 16
150 F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E  chimeric Ad5/Fib16

180 D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y  genbank Fiber 16
180 D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y  chimeric Ad5/Fib16

210 T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S  genbank Fiber 16
210 T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S  chimeric Ad5/Fib16

240 L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A  genbank Fiber 16
240 L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A  chimeric Ad5/Fib16

270 Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F  genbank Fiber 16
270 Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F  chimeric Ad5/Fib16

300 P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E  genbank Fiber 16
300 P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E  chimeric Ad5/Fib16

330 E A P E T T E V T L I T S P F F F S Y I R E D D  genbank Fiber 16
330 E A P E T T E V T L I T S P F F F S Y I R E D D  chimeric Ad5/Fib16
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Decoration 'Decoration #1': Box residues that differ from genbank Fiber 16 .

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